SEQLIST.TXT

SEQUENCE LISTING

```
<110> Novartis AG
      BOLLEKENS, Jacques
      CHIBOUT, Salah-Dine
      VONDERSCHER, Jacky
      LEGAY, Francois
      CORDIER, Andre
      PAPOIAN, Ruben
      SCHERER, Andreas
<120> Use of Fibroblast Growth Factor
  Fragments
<130> 33264-US-PCT
<140> 10578470
<141> 2008-04-18
<150> PCT/EP2004/012572
<151> 2004-11-05
<160> 14
<170> FastSEQ for Windows Version 4.0
<210> 1
<211> 251
<212> PRT
<213> Homo sapiens
<400> 1
Met Leu Gly Ala Arg Leu Arg Leu Trp Val Cys Ala Leu Cys Ser Val
Cys Ser Met Ser Val Leu Arg Ala Tyr Pro Asn Ala Ser Pro Leu Leu
20 25 30
Gly Ser Ser Trp Gly Gly Leu Ile His Leu Tyr Thr Ala Thr Ala Arg
35 40 45
Asn Ser Tyr His Leu Gln Ile His Lys Asn Gly His Val Asp Gly Ala 50 _ 60 _ 60
Pro His Gln Thr Ile Tyr Ser Ala Leu Met Ile Arg Ser Glu Asp Ala 65 70 75 80
Gly Phe Val Val Ile Thr Gly Val Met Ser Arg Arg Tyr Leu Cys Met
                                       90
Asp Phe Arg Gly Asn Ile Phe Gly Ser His Tyr Phe Asp Pro Glu Asn
             100
                                  105
Cys Arg Phe Gln His Gln Thr Leu Glu Asn Gly Tyr Asp Val Tyr His
115 120 125
                              120
Ser Pro Gln Tyr His Phe Leu Val Ser Leu Gly Arg Ala Lys Arg Ala
130 135 140
Phe Leu Pro Gly Met Asn Pro Pro Pro Tyr Ser Gln Phe Leu Ser Arg
                                           155
                     150
                                                                 160
Arg Asn Glu Ile Pro Leu Ile His Phe Asn Thr Pro Ile Pro Arg Arg
                 165
                                       170
His Thr Arg Şer Ala Glu Asp Asp Şer Glu Arg Asp Pro Leu Asn Val
             180
                                                        190
                                  185
Leu Lys Pro Arg Ala Arg Met Thr Pro Ala Pro Ala Ser Cys Ser Gln
                              200
                                                    205
Glu Leu Pro Ser Ala Glu Asp Asn Ser Pro Met Ala Ser Asp Pro Leu
    210
                          215
                                               220
Gly Val Val Arg Gly Gly Arg Val Asn Thr His Ala Gly Gly Thr Gly 225 230 235
```

Pro Glu Gly Cys Arg Pro Phe Ala Lys Phe Ile 245 250

```
<210> 2
<211> 75
<212> PRT
<213> Homo sapiens
<400> 2
His Thr Arg Ser Ala Glu Asp Asp Ser Glu Arg Asp Pro Leu Asn Val
                                       10
Leu Lys Pro Arg Ala Arg Met Thr Pro Ala Pro Ala Ser Cys Ser Gln
                                   25
             20
Glu Leu Pro Ser Ala Glu Asp Asn Ser Pro Met Ala Ser Asp Pro Leu
                              40
   Val Val Arg Gly Gly Arg Val Asn Thr His Ala Gly Gly Thr Gly 50 60
Pro Glu Gly Cys Arg Pro Phe Ala Lys Phe Ile
<210> 3
<211> 756
<212> DNA
<213> Homo sapiens
<400> 3
atgttggggg cccgcctcag gctctgggtc tgtgccttgt gcagcgtctg cagcatgagc 60
gtcctcagag cctatcccaa tgcctcccca ctgctcggct ccagctgggg tggcctgatc
                                                                         120
cacctgtaca cagccacagc caggaacagc taccacctgc agatccacaa gaatggccat
                                                                        180
gtggałggcg caccccatca gaccatctac agtgccctga tgatcagatc agaggatgct 240
ggctttgtgg tgattacagg tgtgatgagc agaagatacc tctgcatgga tttcagaggc 300
aacatttttg gatcacacta tttcgacccg gagaactgca ggttccaaca ccagacgctg 360
gaaaacgggt acgacgtcta ccactctcct cagtatcact tcctggtcag tctgggccgg 420
gcgaagagag ccttcctgcc aggcatgaac ccaccccgt actcccagtt cctgtcccgg 480
aggaacgaga tccccctaat tcacttcaac acccccatac cacggcggca cacccggagc 540
gccgaggacg actcggagcg ggaccccctg aacgtgctga agccccgggc ccggatgacc 600
ccggccccgg cctcctgttc acaggagctc ccgagcgccg aggacaacag cccgatggcc 660 agtgacccat taggggtggt caggggcggt cgagtgaaca cgcacgctgg gggaacgggc 720
                                                                         756
ccggaaggct gccgccctt cgccaagttc atctag
<210> 4
<211> 228
<212> DNA
<213> Homo sapiens
<400> 4
cacacccgga gcgccgagga cgactcggag cgggaccccc tgaacgtgct gaagccccgg 60
gcccggatga ccccggcccc ggcctcctgt tcacaggagc tcccgagcgc cgaggacaac 120\,
agcccgatgg ccagtgaccc attaggggtg gtcaggggcg gtcgagtgaa cacgcacgct 180 ggggggaacgg gcccggaagg ctgccgccc ttcgccaagt tcatctag 228
<210> 5
<211> 79
<212> PRT
<213> Homo sapiens
<400> 5
Gln Arg Asp Pro Val Gly Arg Tyr Glu Pro Ala Gly Gly Asp Ala Asn
1
            Arg Pro Gly Gly Ser Tyr Pro Ala Ala Ala Ala Lys 20 25 30
```

Val Tyr Ser Leu Phe Arg Glu Gln Asp Ala Pro Val Ala Gly Leu Gln

```
SEQLIST.TXT
45
```

Pro Val Glu Arg Ala Gln Pro Gly Trp Gly Ser Pro Arg Arg Pro Thr 50 55 60
Glu Ala Glu Ala Arg Arg Pro Ser Arg Ala Gln Gln Ser Arg Arg 65 70 75

<210> 6 <211> 79

<212> PRT <213> Mus musculus

...

<210> 7 <211> 63 <212> PRT <213> Homo sapiens

 $<\!400\!>$ 7 Ala Thr Leu Gly Gly Pro Glu Glu Glu Ser Thr Ile Glu Asn Tyr Ala 1 5 10 15 Ser Arg Pro Glu Ala Phe Lys Ala Asp Glu Phe Leu Asn Trp His Ala 20 25 30

Leu Phe Glu Ser Ile Lys Arg Lys Leu Pro Phe Leu Asn Trp Asp Ala

Phe Pro Lys Leu Lys Gly Leu Arg Ser Ala Thr Pro Asp Ala Gln
50 55 60

<210> 8 <211> 77 <212> PRT <213> Homo sapiens

<210> 9 <211> 76 <212> PRT <213> Homo sapiens

```
SEQLIST.TXT
<400> 9
Ala Thr Leu Gly Gly Pro Glu Glu Glu Ser Thr Ile Glu Asn Tyr Ala
1 5 10 15
Ser Arg Pro Glu Ala Phe Asn Thr Pro Phe Leu Asn Ile Asp Lys Leu 20 25 30
Arg Ala Phe Lys Ala Asp Glu Phe Leu Asn Met His Ala Leu Phe Glu 40 45
Ser Ile Lys Arg Lys Leu Pro Phe Leu Asn Trp Asp Ala Phe Pro Lys 50 _ 60
Leu Lys Gly Leu Arg Ser Ala Thr Pro Asp Ala Gln 65 70 75
<210> 10
<211> 49
<212> PRT
<213> Homo sapiens
Cys Ala Tyr Thr Phe Leu Val Pro Glu Gln Arg Ile Thr Gly Pro Ile
<400> 10
Cys Val Asn Thr Lys Gly Gln Asp Ala Ser Thr Ile Lys Asp Met Ile
                                   25
Thr Arg Met Asp Leu Glu Asn Leu Lys Asp Val Leu Ser Arg Gln Lys 35 40 45
Arg
<210> 11
<211> 30
<212> PRT
<213> Mus musculus
<400> 11
Cys Ser Tyr Thr Phe Leu Val Pro Glu Gln Lys Ile Thr Gly Pro Ile
1 5 10 15
```

Cys Val Asn Thr Lys Gly Gln Asp Ala Gly Thr Ile Lys Asp 20 25 30

<210> 12 <211> 20 <212> PRT <213> Mus musculus

<400> 12 Met Ile Thr Arg Met Asp Leu Glu Asn Leu Lys Asp Val Leu Ser Arg 10 Gln Lys Arg Glu 20

<210> 13 <211> 24 <212> PRT <213> Homo sapiens

<400> 13 Ser Thr Ile Lys Asp Met Ile Thr Arg Met Asp Leu Glu Asn Leu Lys
1 10 15 Asp Val Leu Ser Arg Gln Lys Arg 20

SEQLIST.TXT